
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Sep 14 16:03:55 EDT 2007

Validated By CRFValidator v 1.0.3

Application No: 10590633 Version No: 1.0

Input Set:

Output Set:

Started: 2007-09-04 12:29:12.260 **Finished:** 2007-09-04 12:29:12.868

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 608 ms

Total Warnings: 5

Total Errors: 2

No. of SeqIDs Defined: 6

Actual SeqID Count: 6

Error code		Error Description			
W	213	Artificial or Unknown found in <213> in SEQ ID (1)			
W	213	Artificial or Unknown found in <213> in SEQ ID (3)			
W	213	Artificial or Unknown found in <213> in SEQ ID (4)			
E	257	Invalid sequence data feature in <221> in SEQ ID (4)			
E	257	Invalid sequence data feature in <221> in SEQ ID (4)			
W	213	Artificial or Unknown found in <213> in SEQ ID (5)			
W	213	Artificial or Unknown found in <213> in SEQ ID (6)			

SEQUENCE LISTING

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<110> ZACHARIAS, DAVID ALAN
<120> FUNCTIONAL GENOMICS AND GENE TRAPPING IN HAPLOID OR
      HYPODIPLOID CELLS
<130> UFRF1100-1(055932-0203)
<140> 10590633
<141> 2007-09-04
<150> PCT/US2005/06309
<151> 2005-02-25
<150> 60/548,509
<151> 2004-02-26
<160> 6
<170> PatentIn Ver. 3.3
<210> 1
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1
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                                     10
Gln Lys
<210> 2
<211> 17
<212> DNA
<213> Mus musculus
<400> 2
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<210> 3
<211> 68
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Consensus
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ccgggctt
<210> 4
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Consensus
     motif
<220>
<221> MOD_RES
<222> (3)..(5)
<223> Variable amino acid
<220>
<221> MOD_RES
<222> (6)
<223> Ser or Thr
<400> 4
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                5
<210> 5
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Illustrative
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<210> 6
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: C-term tail
      peptide of GFP: Yck2p
<400> 6
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 1
                  5
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<400> 3